

FIG. 1

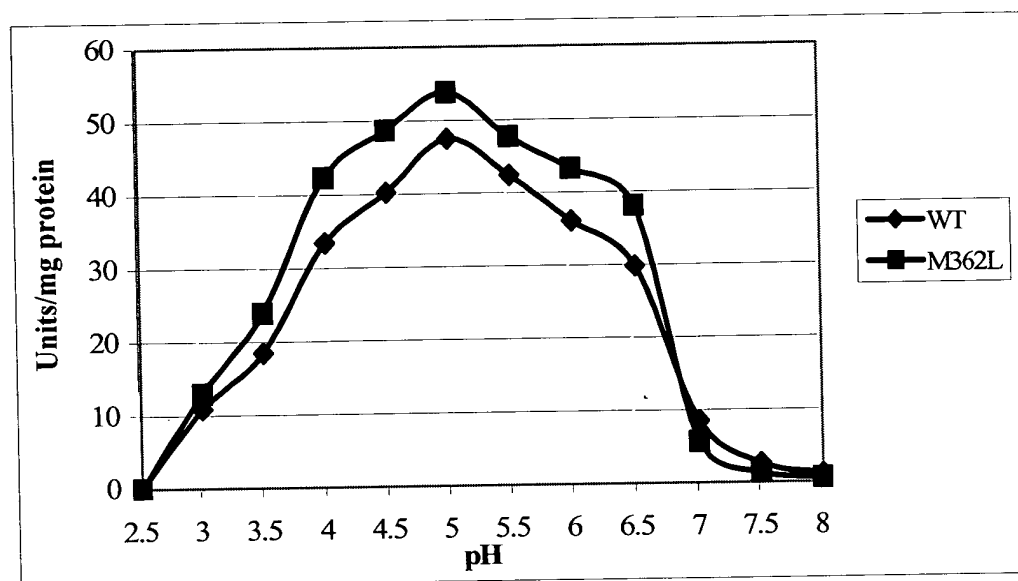


FIG. 2

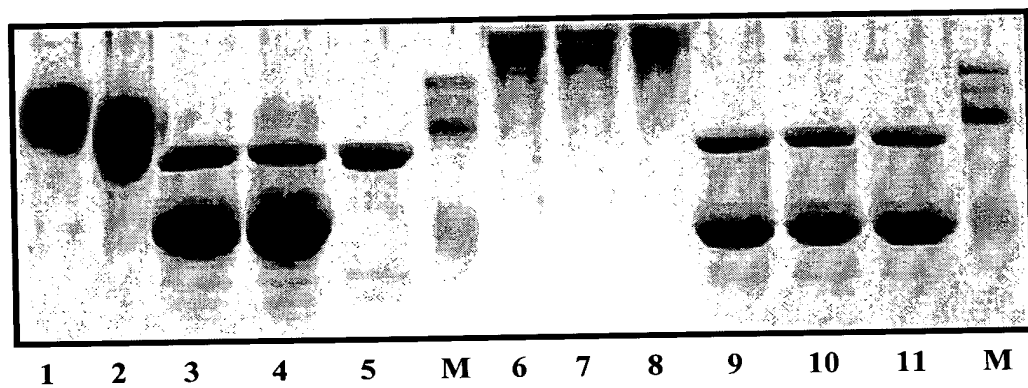


FIG. 3

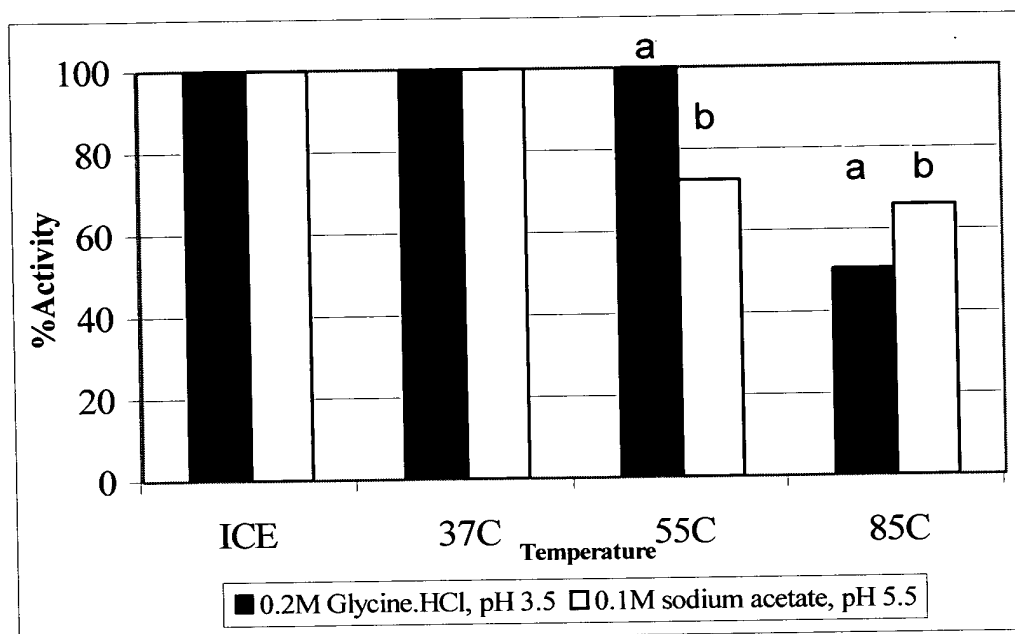


FIG. 4

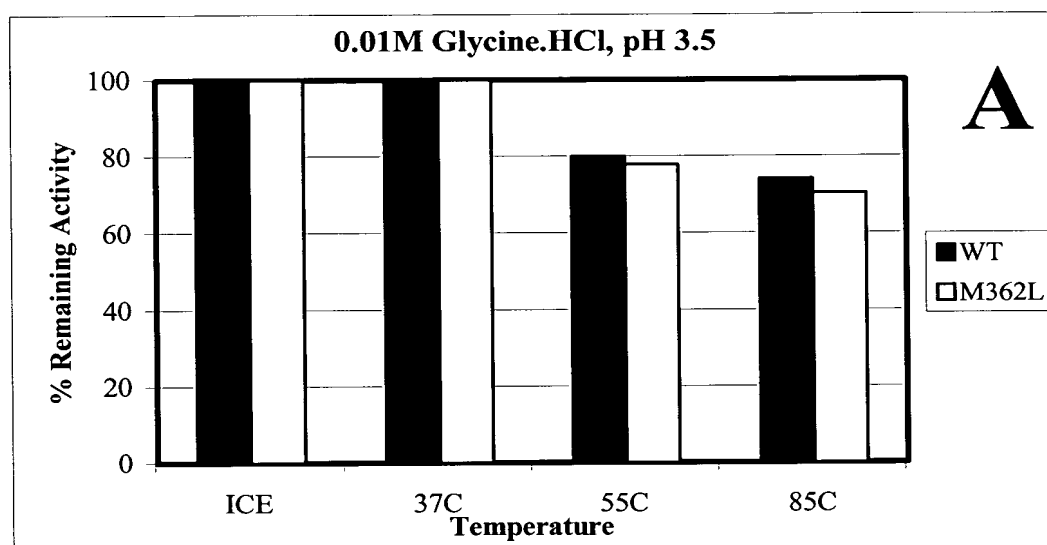


FIG. 5A

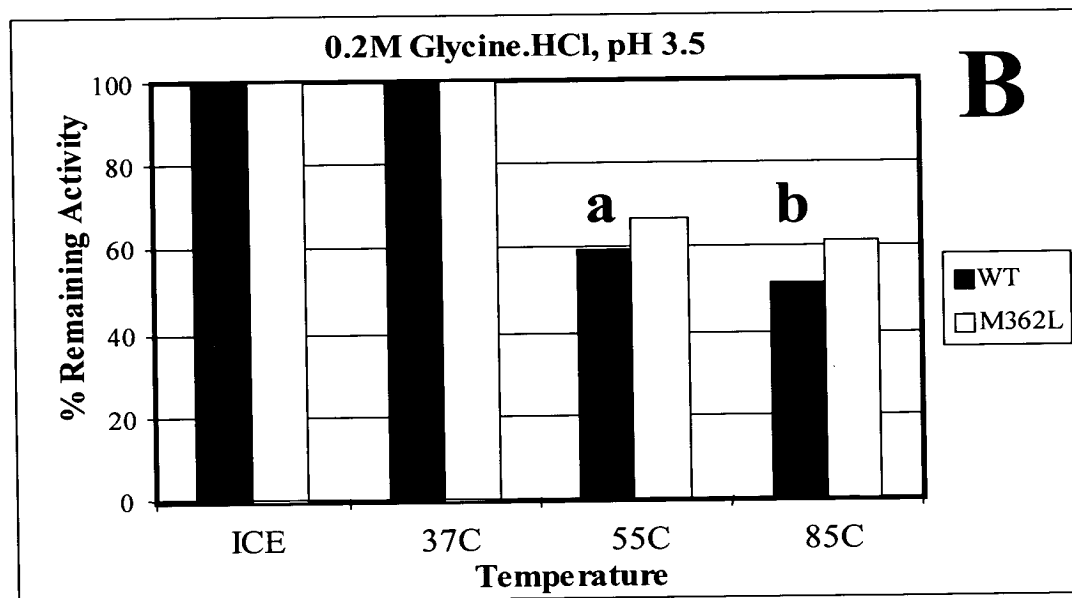


FIG. 5B

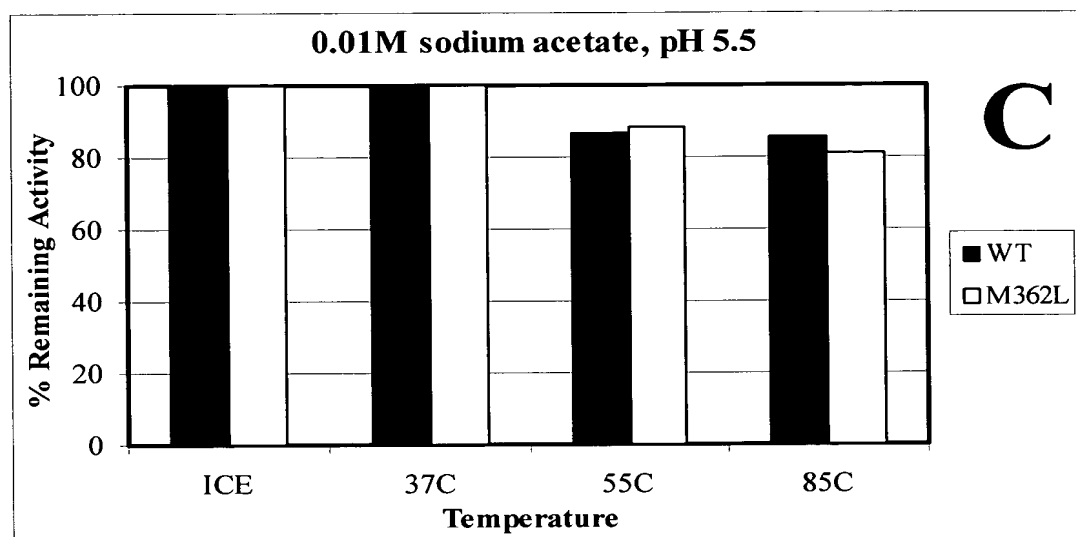


FIG. 5C

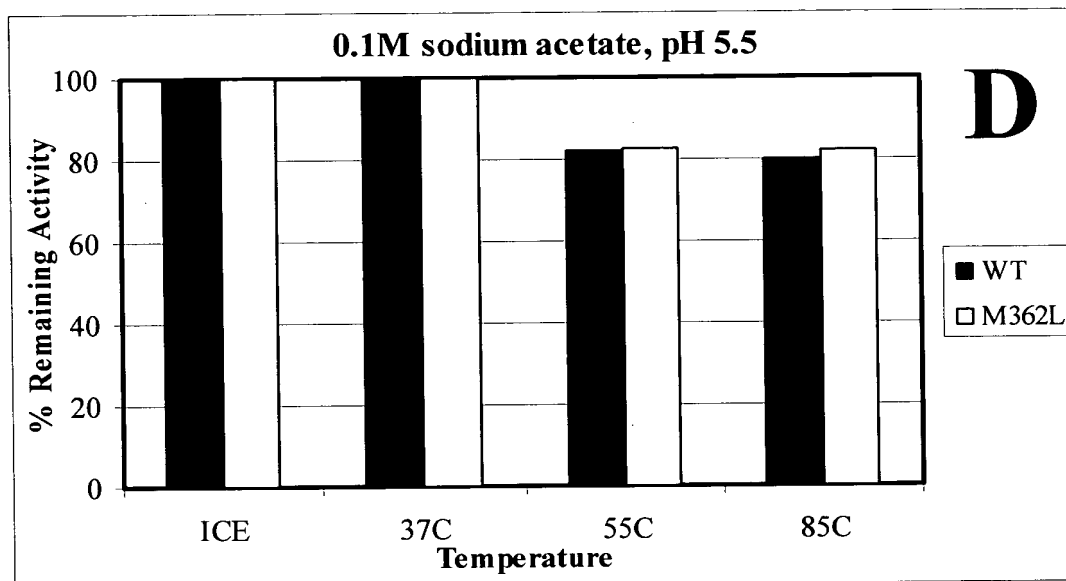


FIG. 5D

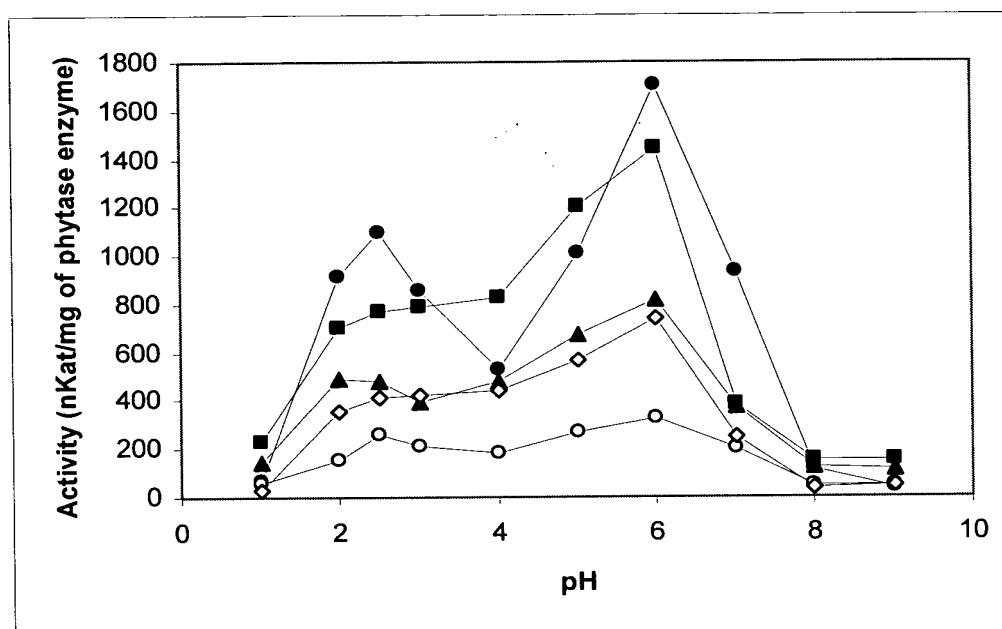


FIG. 6

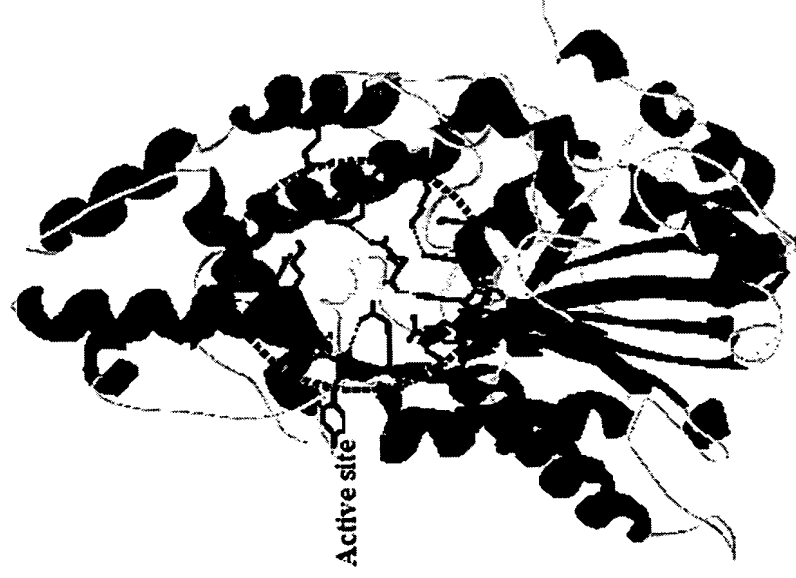


FIG. 7A

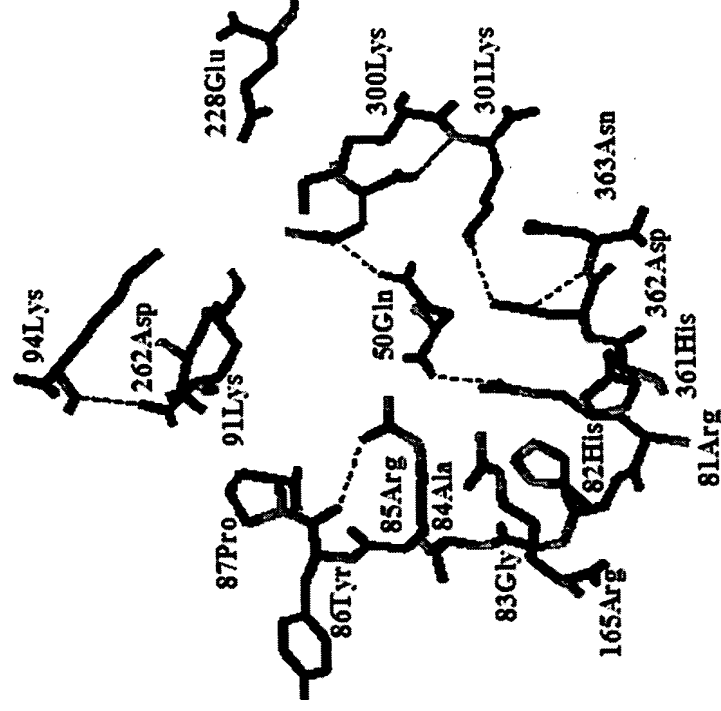


FIG. 7B

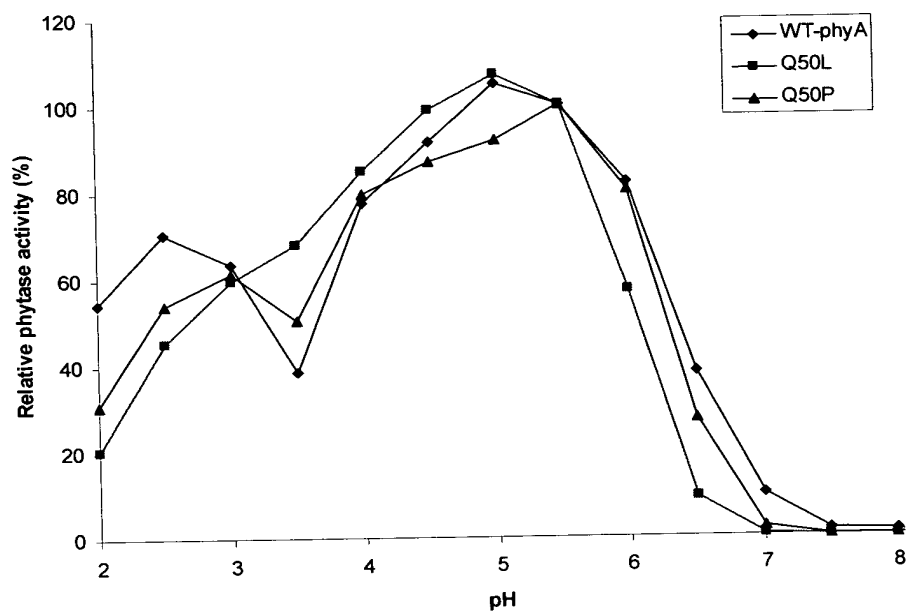


FIG. 8A

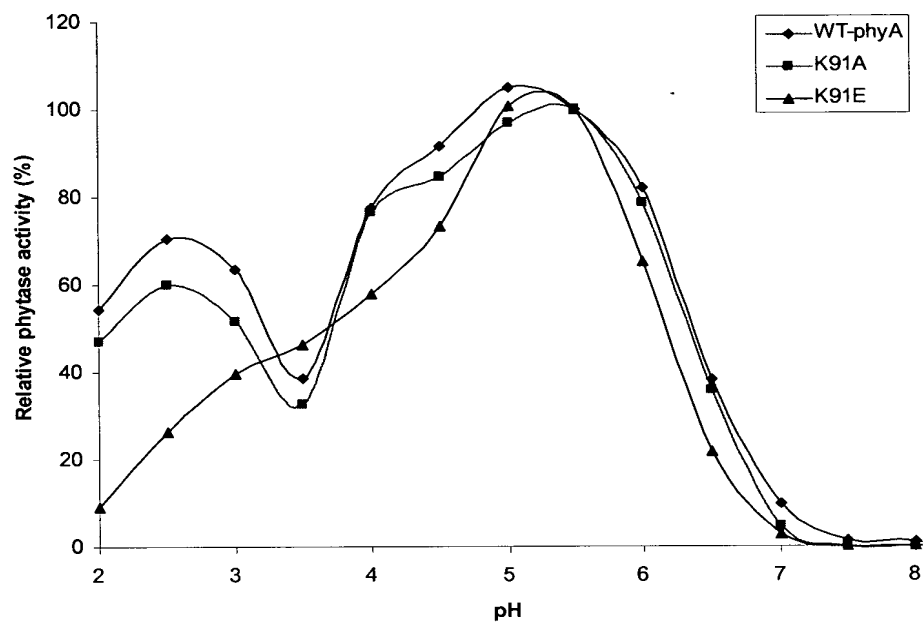


FIG. 8B

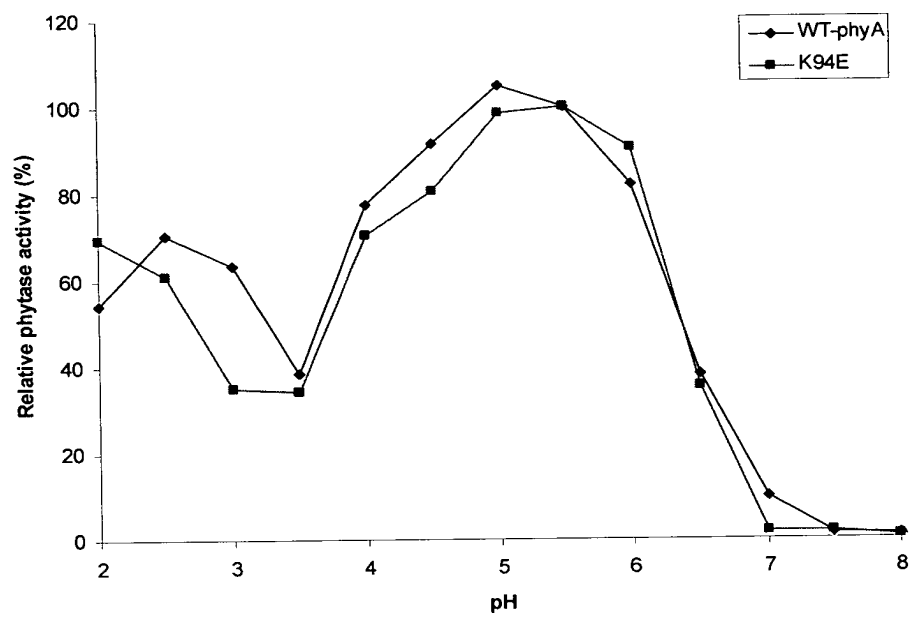


FIG. 8C

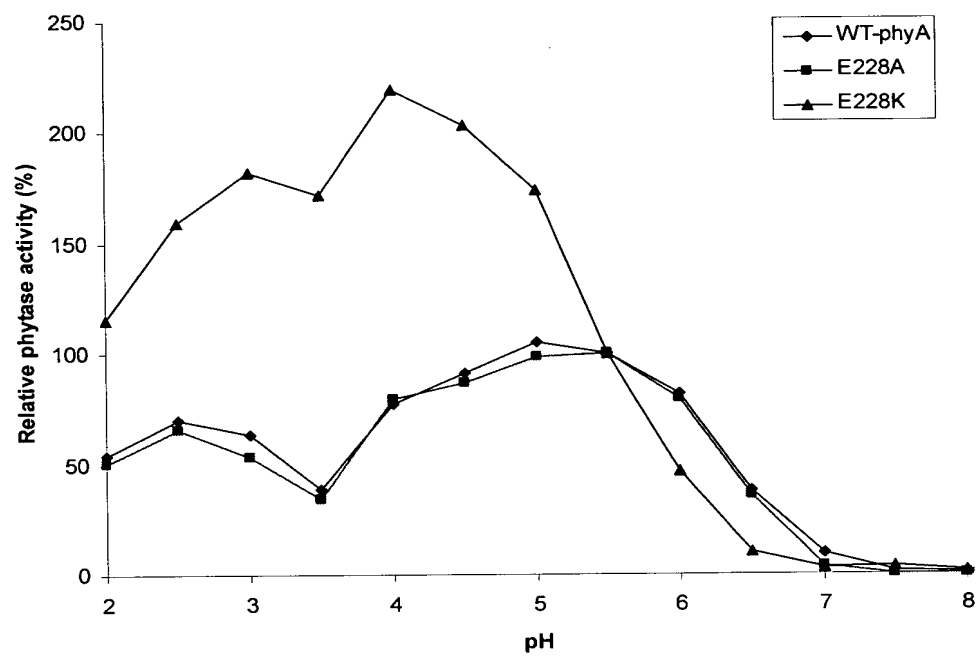


FIG. 8D

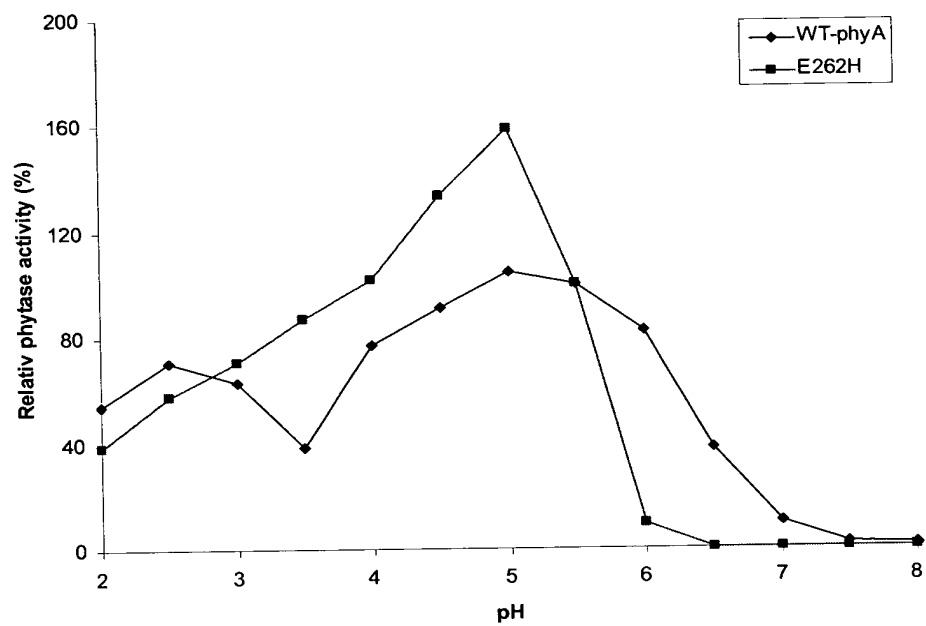


FIG. 8E

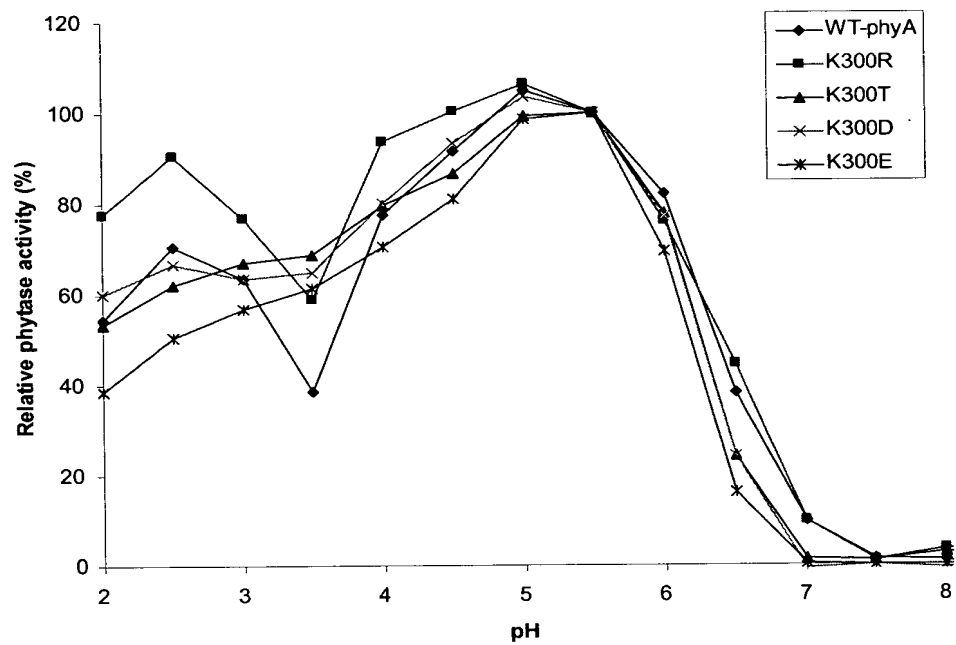


FIG. 8F

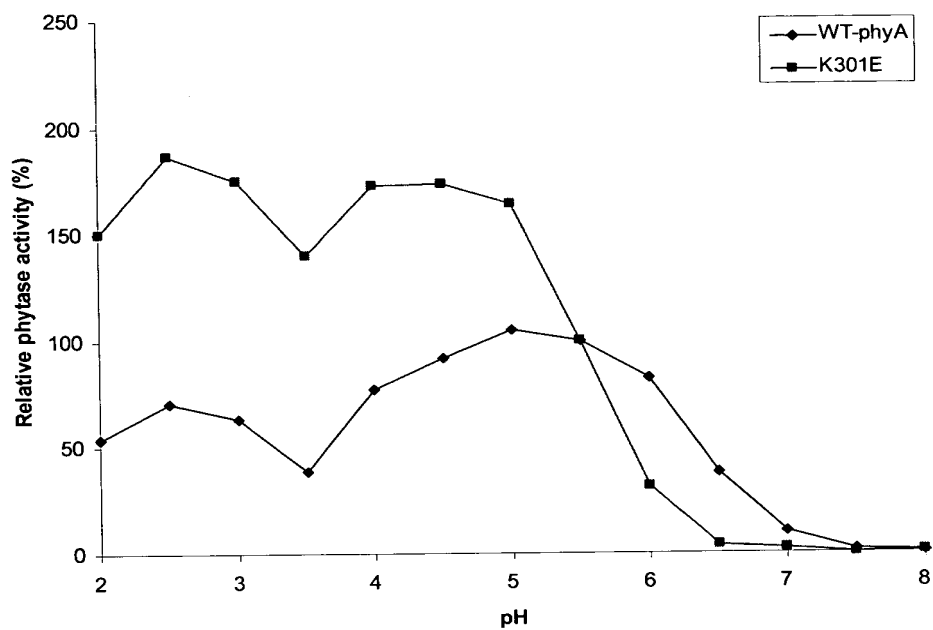


FIG. 8G

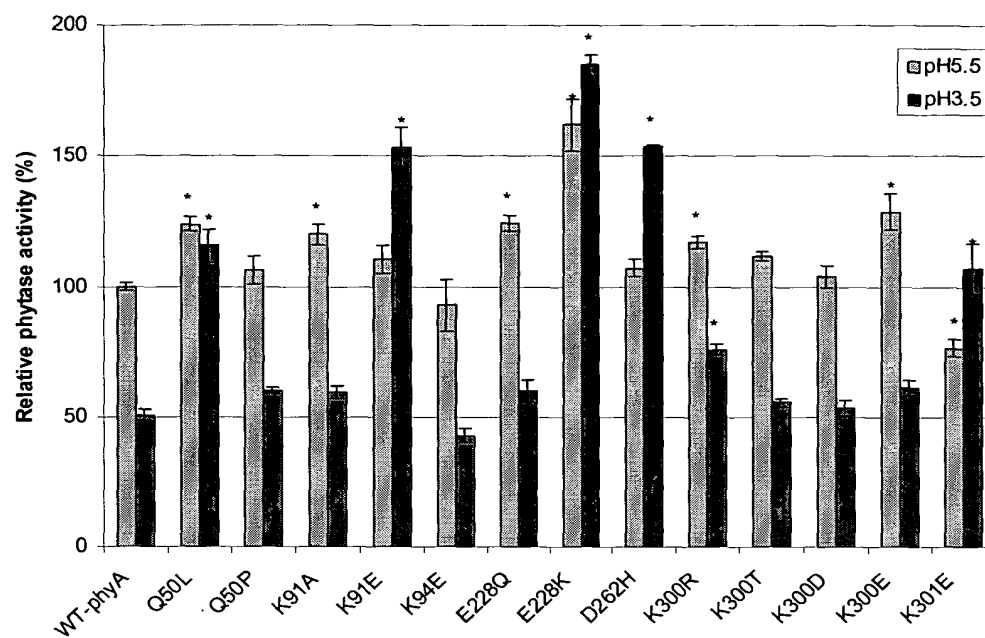


FIG. 9

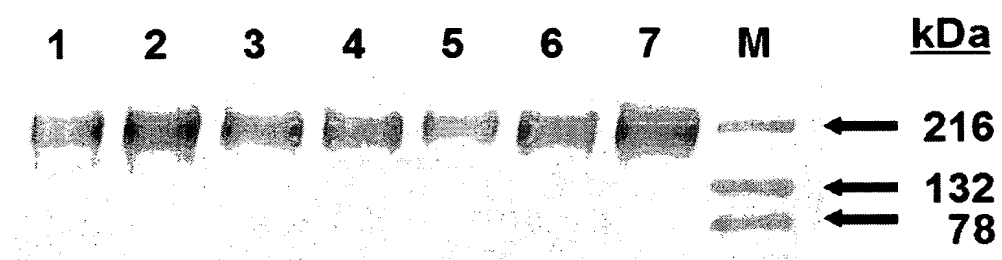


FIG. 10A



FIG. 10B

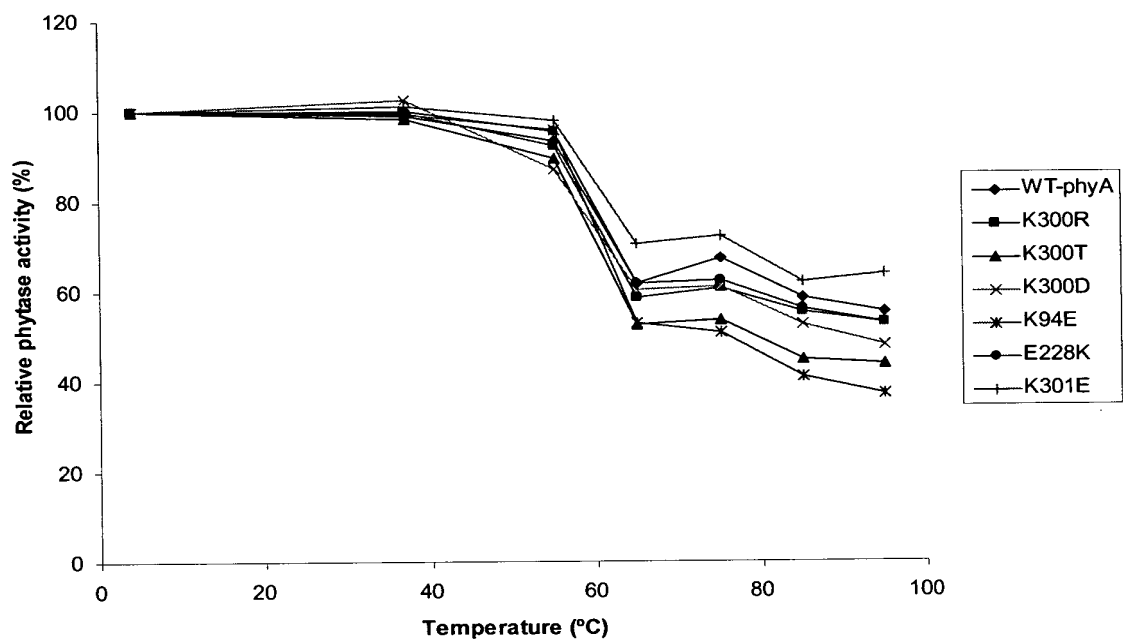


FIG. 11

1st line: Asp. terreus phytase
 2nd line: Asp. niger phytase
 3rd line: Asp. fumigatus phytase

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1  MGVFVLLSI  ATLFGSTSGT  ALGPRGNHSD  CTSVDRGYQC  EPESHKWGL
1  MGVSALLPL  YLLSGVTSGL  AVPASRNQSS  CDTVDQGYQC  FSETSHLWQ
1  MVTLTFLLSA  AYLLSGRVS-  AAPSSAGSKS  CDTVDLGYQC  SPATSHLWQ
****                               SSS  SSSS

51  YAPYFSLQDE  SPFPLDVDD  CHITFVOVLA  RHGARSPTDS  KTKAAYAATIA
51  YAPFFSLANE  SVISPEVPAG  CRVITFAOVLS  RHGARYPTDS  KGKKYSALIE
50  YSPFFSLEDE  LSVSSKLEKD  CRITLVOVLS  RHGARYPTSS  KSKKYKKIVT
****          SSS          S  SSSSSSSSSSSSS SSS  HH  HHHHHHHHHH

101  ALOKNATALP  GKYAFLKSYN  YSMGSENINP  FGRNQLQDLG  AOFYRRYDTL
101  ELQONATTED  GKYAFLKTYN  YSLGADDLTP  FGEQELVNSG  IKFYQRYESL
100  ALOANATDEK  GKFAFLKTYN  YTLGADDLTP  FGEQQLVNSG  IKFYQRYKAL
****  HHHHHH          HHHH          SSSH  HHHHHHHHHH  HHHHH  HH

151  TRHINPFVRA  ADSSRVHESA  EKEVEGEFQA  RQGDPEHANEH  QPSPRVDVVI
151  TRNIVPFIRS  SGSSRVIASG  KKFIEGFQST  KIKDERAQPG  QSSPKTDVVI
150  ARSVVPFIRA  SGSDRVIASG  EKFIIEGFQQA  KLADEGA- TN  RAAPATSMII
****  H          SSSS  SS  HHHHHHHH  HHHHHHHHHH  HH          SSS

201  PEGTAYNNTL  EHSICTAFEA  STVGDAADN  FTAVEAPATA  KRLEADLPV
201  SEASSSNNTL  DPGTCTVFED  SELADTVEAN  FTATEVESIR  QRLNDLSGV
199  PESETFNNTL  DHGVCTKFEA  SOLGDEVAAN  FTALFAPDIR  ARAEKHLPGV
****          HHHH          HHHHHHHH  HHHH  HHHHHH

251  QLSADDVNL  MAMCPPEIVS  LTDDAHTLSP  FCDLFTAAEW  TOYNYILSLD
251  TLTDEVTYL  MDMCSFDTIS  TSTVDTKLSP  FCDLFTHDEW  INYDYLOSLK
249  TLTDEDVSL  MDMCSFDTVA  RTSDASQLSP  FCQLFTHNEW  KKYNLYQSLG
****          HHHHHH  HHHHHHHH  HHH  HHHH  HHHHHHHHHH

301  KYYGYGGNP  LGFVOGVGWA  NELIARLTRS  PVHDHTCVNN  TLDANPATFP
301  KYYGHGAGNP  LGPTQGVGYA  NELIARLTHS  PVHDDTSSNH  TLDSSPATFP
299  KYYGYGAGNP  LGPAQGIGFT  NELIARLTRS  PVODHTSTNS  TLVSNPATFP
****  H          HH  HHH  HHHHHHHH          H  HHH

351  LNATLYADFS  HDSNLVSIWF  ALGLYNGTKP  LSOTTVEDIT  RTDGYAAAWT
351  LNSTLYADFS  HDNGIISILE  ALGLYNGTKP  LSTTTVENIT  QTDGFSSAWT
349  LNATMYVDFS  HDNSMVSTFE  ALGLYNGTEE  LSRTSVESAK  ELDGYSASW
****          SSSSSS  HHHHHHHH  H          HHHH

401  VPFAARAYIE  MMOCRAEKOP  LVRVLVNDRV  MPLHGCAVDN  LGRCRKRDDFV
401  VPFAARLYVE  MMOCQAEQEP  LVRVLVNDRV  VPLHGCPVDA  LGRCRTRDSFV
399  VPFGARAYFE  TMOCKSEKEP  LVRALINDRV  VPLHGCDVDK  LGRCRKLNDFV
****          SSSSSS  SSSSS  S  SSSSSSS  SS  S          SSSHHH

451  EGLSFARAGG  NWAECF-
451  RGLSFARSGG  DWAECEA
449  KGLSWARSGG  NWGECEA
****  H          HHH          HHGTT

```

*Red letter shows the mutation site for substrate binding site.
 *Bold letters are known as critical catalytic active sites.

FIG. 12

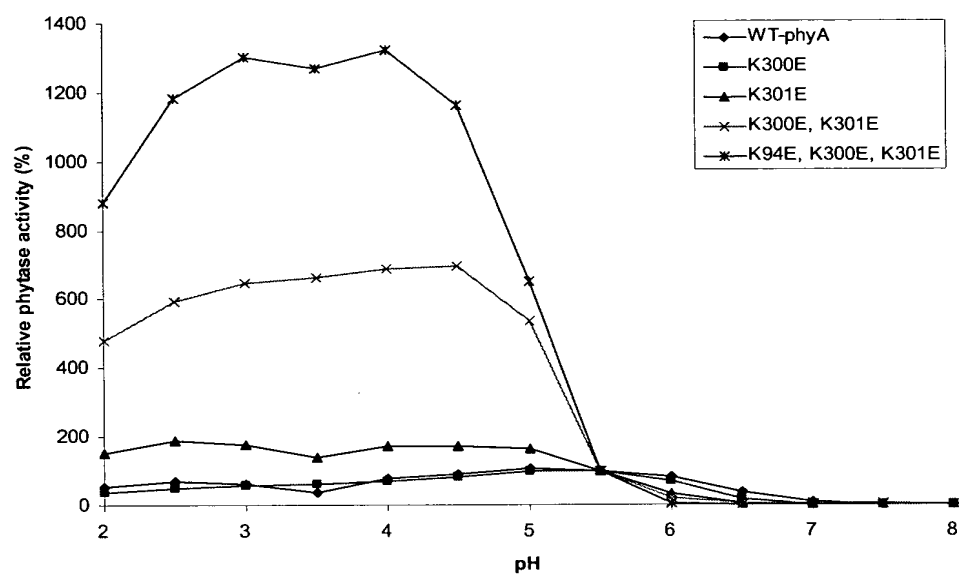


FIG. 13A

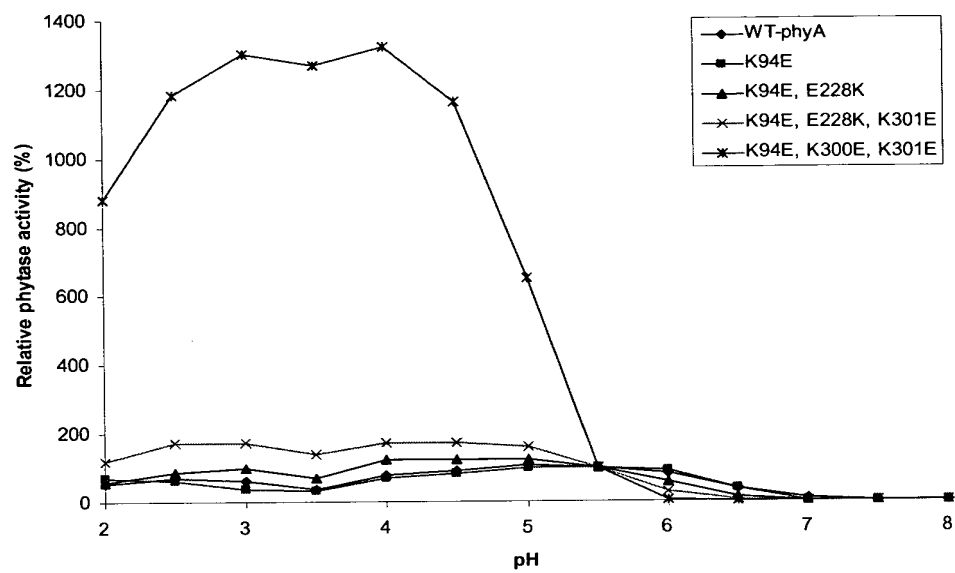


FIG. 13B

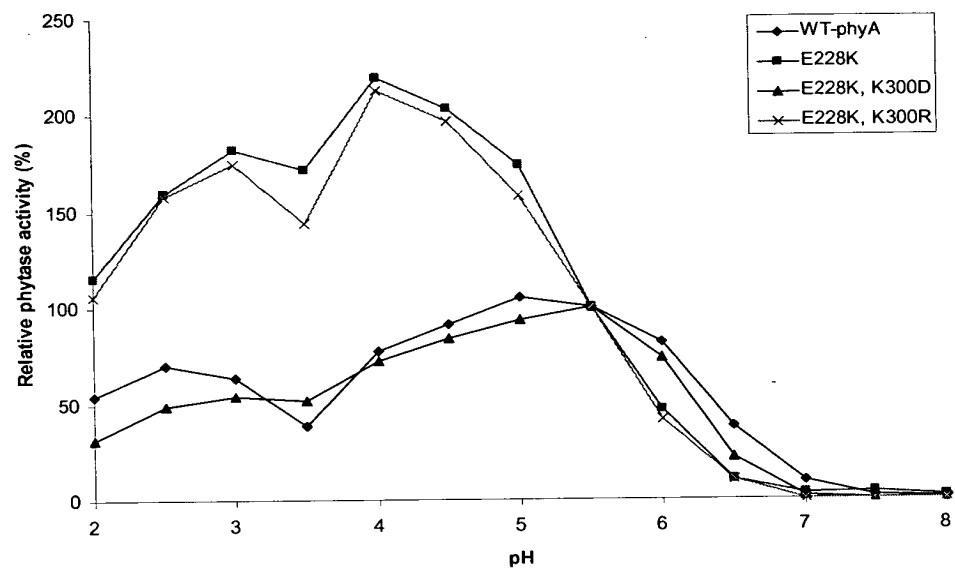


FIG. 13C

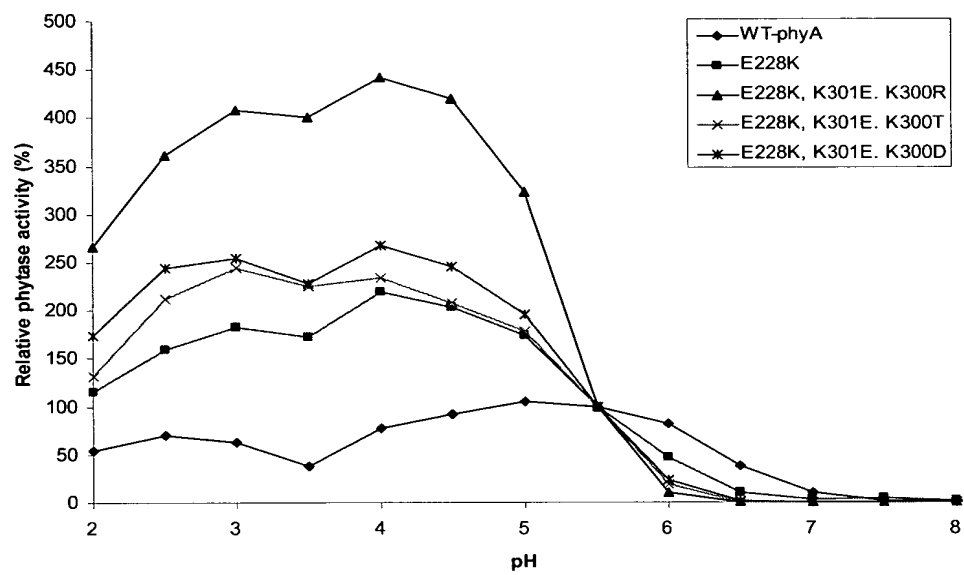


FIG. 13D

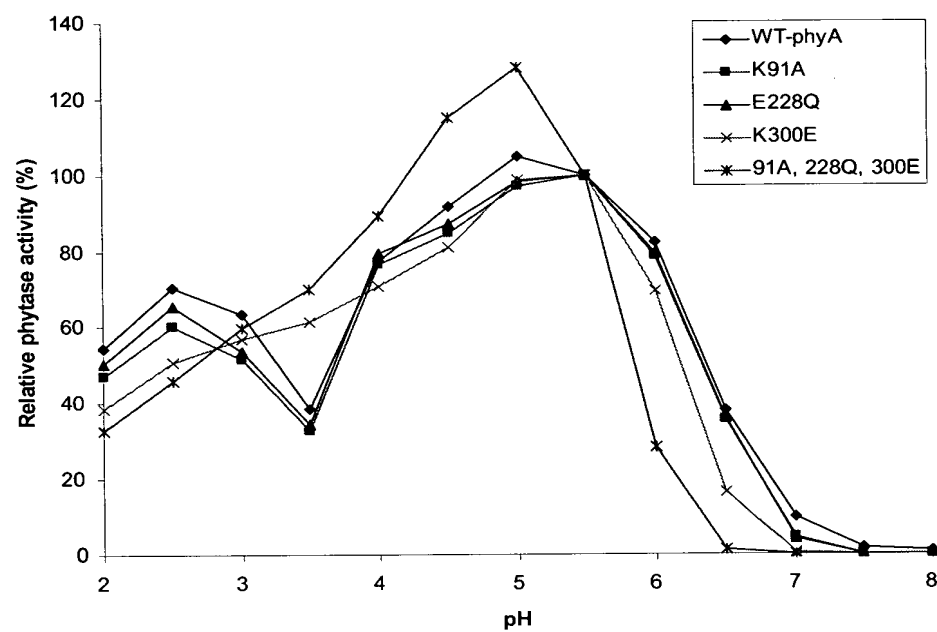


FIG. 13E

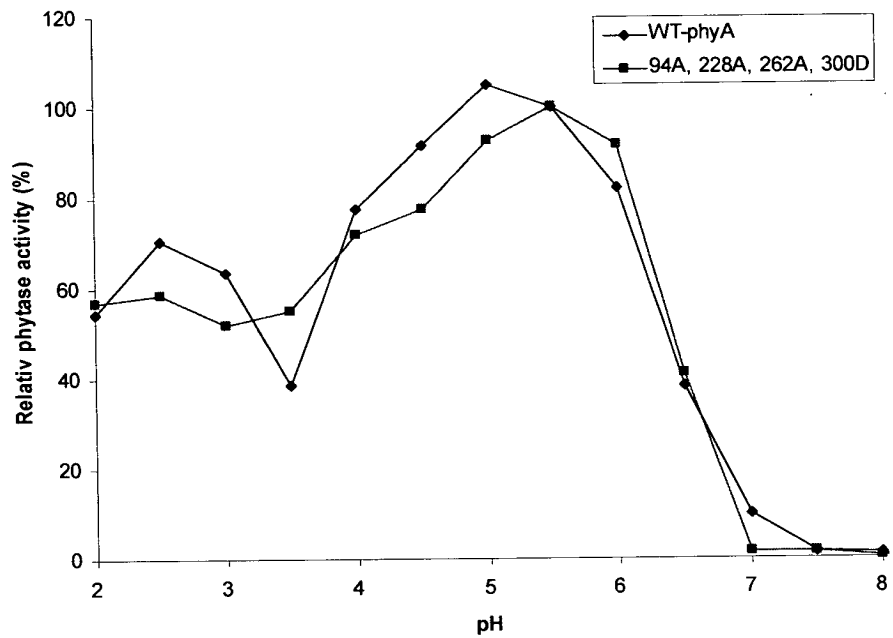


FIG. 13F

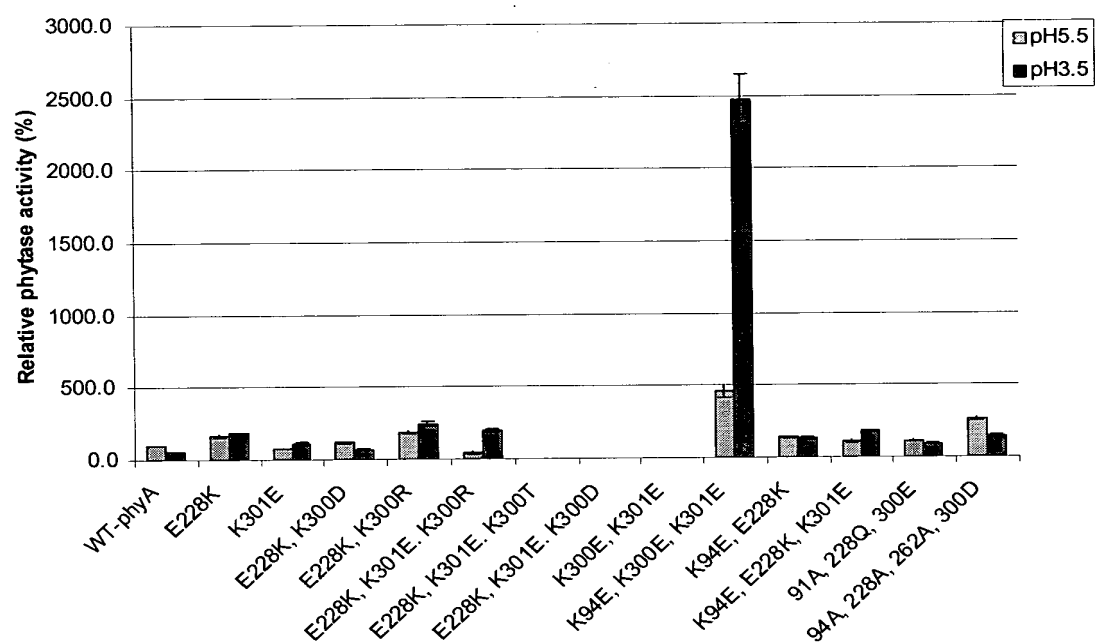


FIG. 14A

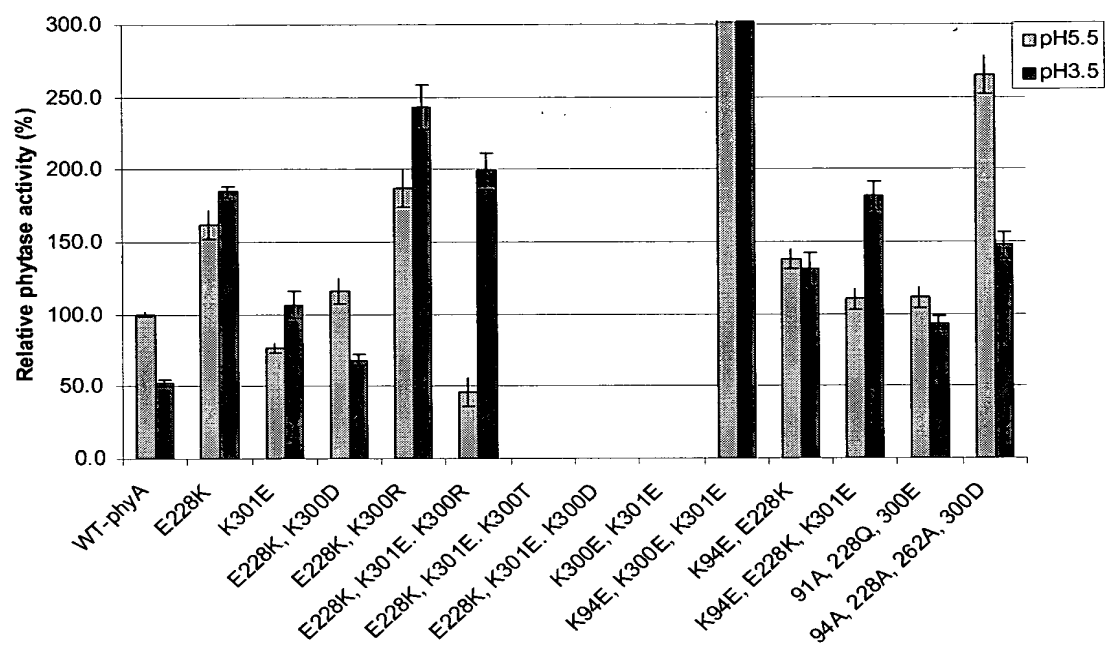


FIG. 14B

Plasma Inorganic Phosphate (PIP) of Pigs Fed Low-P Diets

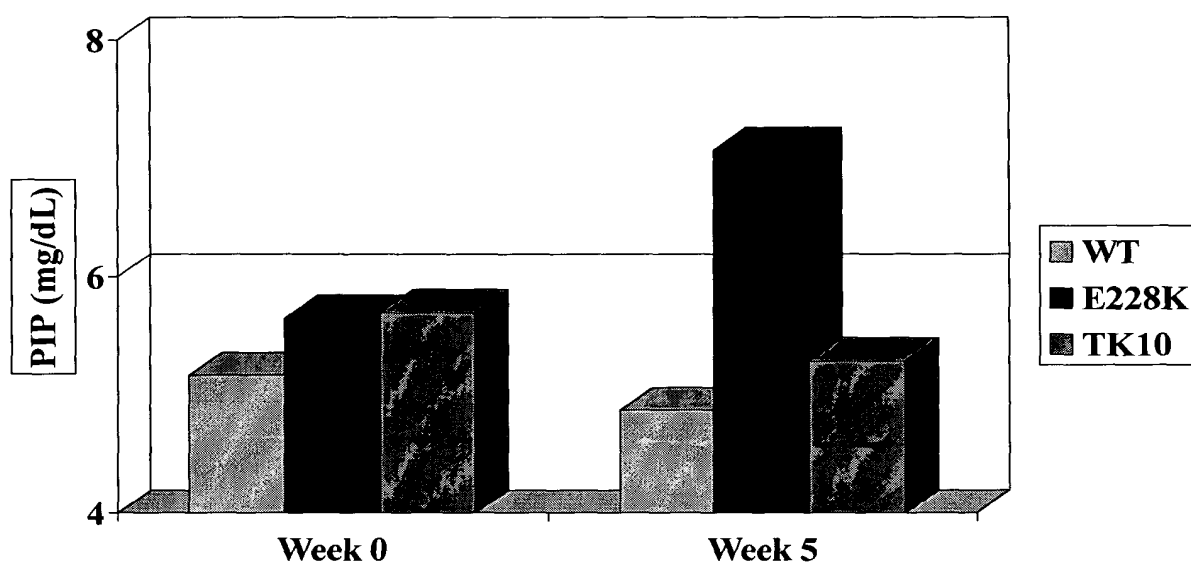


FIG. 15

Plasma Alkaline Phosphatase (AKP) Activity of Pigs Fed Low-P Diets

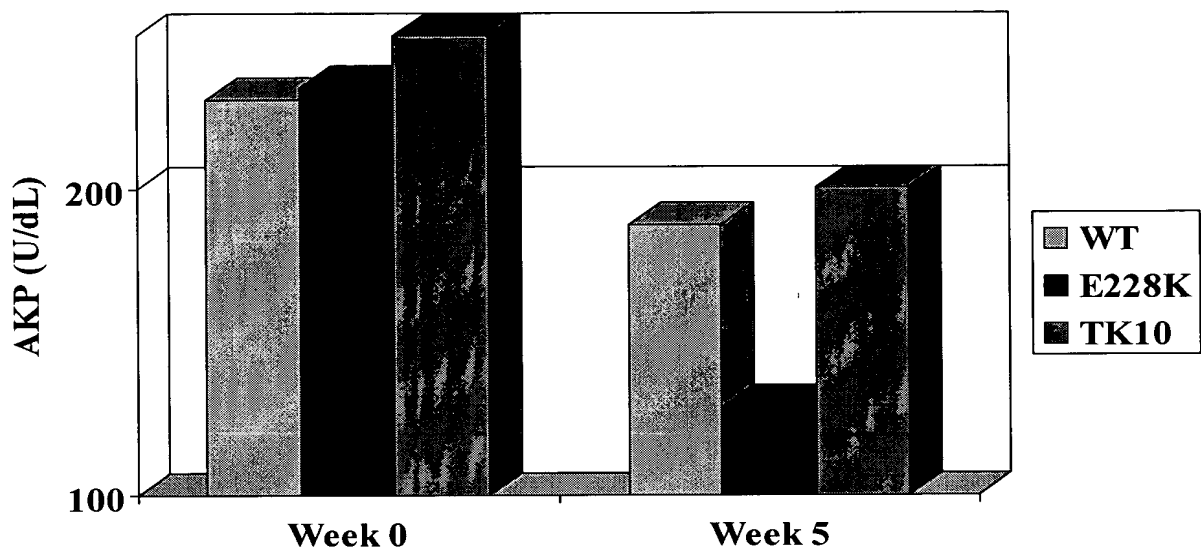


FIG. 16

Average Daily Gain (ADG) of Pigs Fed Low-P Diets

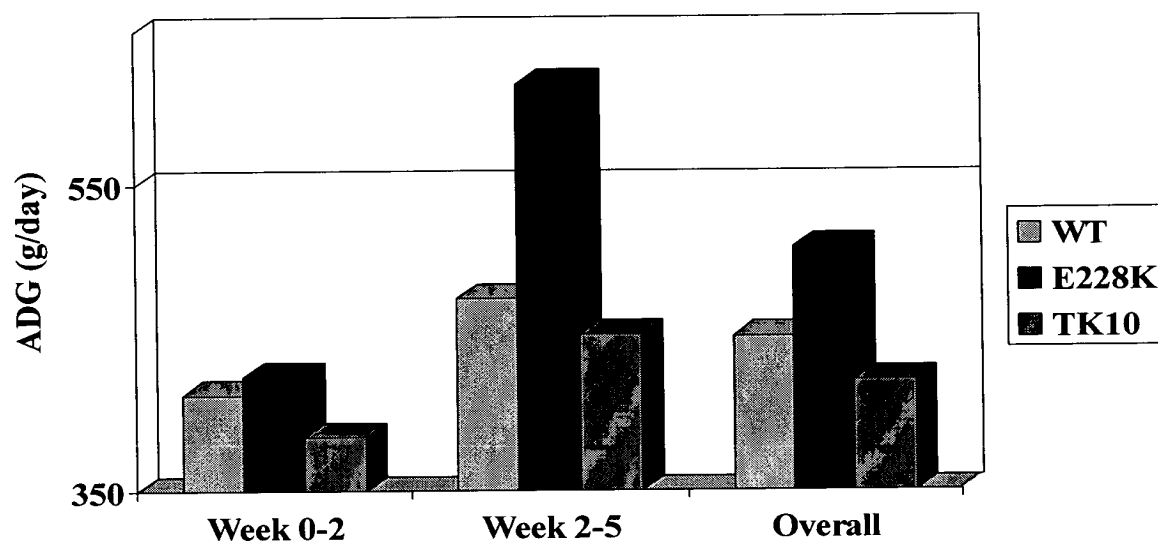


FIG. 17

Gain/Feed of Pigs Fed Low-P Diets

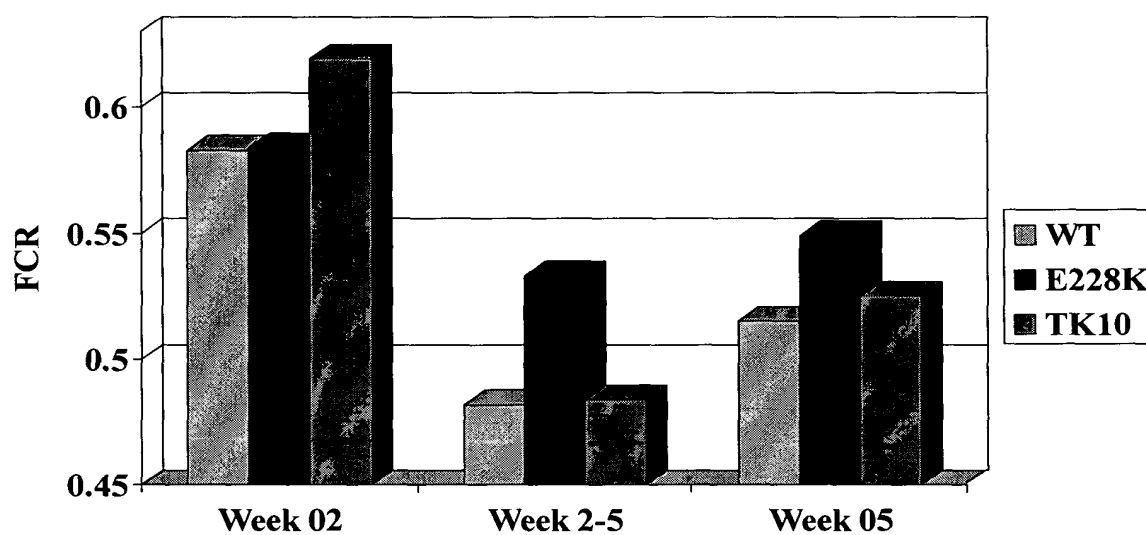


FIG. 18